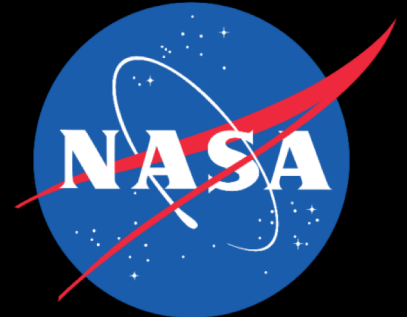
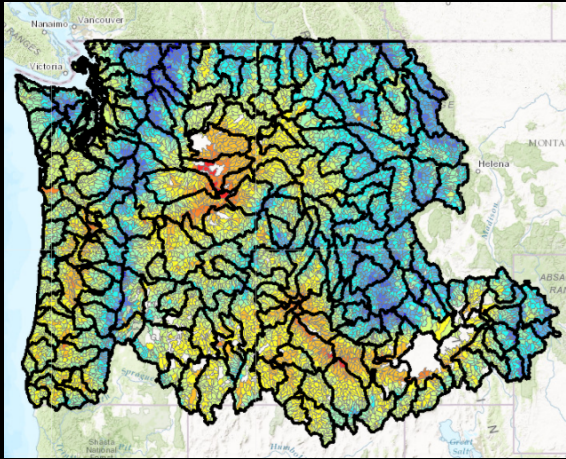


Projecting Effects of Climate Change on River Habitats and Salmonid Populations:

Integrating Remote Sensing, Genomics, and Demography
to Inform Conservation

Brian Hand, Diane Whited, Alisa Wade, Nick Gayeski (Wild Fish
Conservancy), Gordon Luikart (PI)

April 24th, 2018



Urgent need for genomic monitoring and climate change assessments of salmonids

Urgent Need

- Salmonids are “canaries of climate change”
- Multi-millions spent annually on salmonid conservation
- Prioritization of populations for management needed

Clean, Connected, **COLD**
Habitat



Spawning bull trout

Vulnerability to future climate depends on climate exposure, sensitivity & adaptive capacity.

Integrate as key elements of VULNERABILITY

Exposure

**Adaptive
Capacity – rapid
response**

**Sensitivity –
physiological
response**

**Habitat
(Remotely
Sensed)**

Climate

Physical habitat
characteristics

Genomics

Number of
Effective Breeders
Genetic diversity

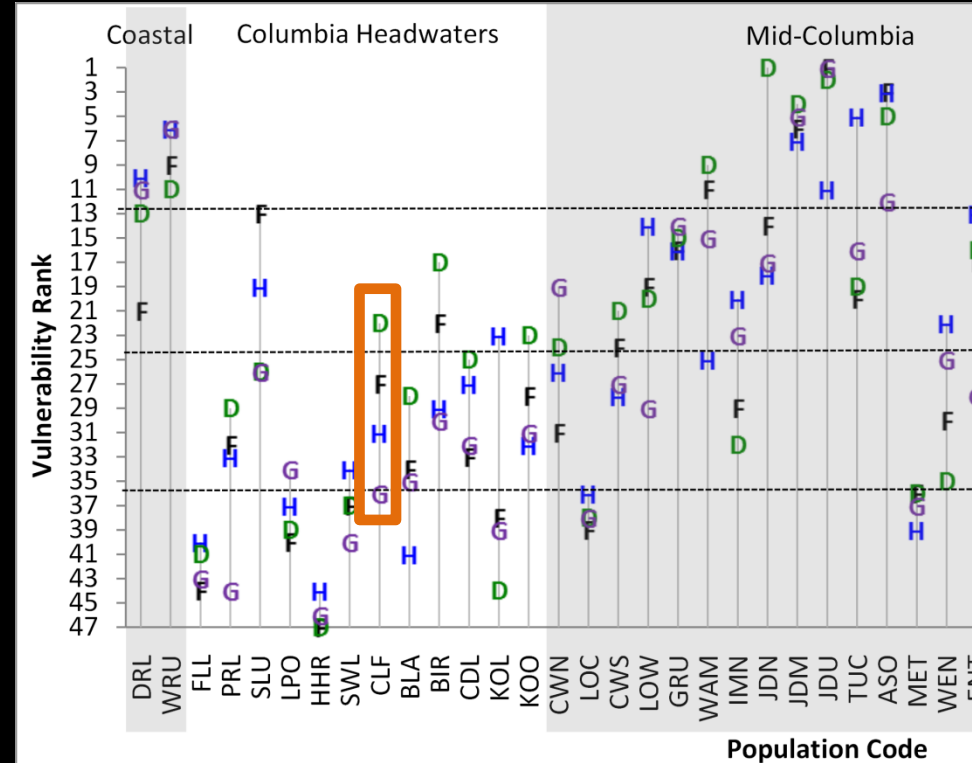
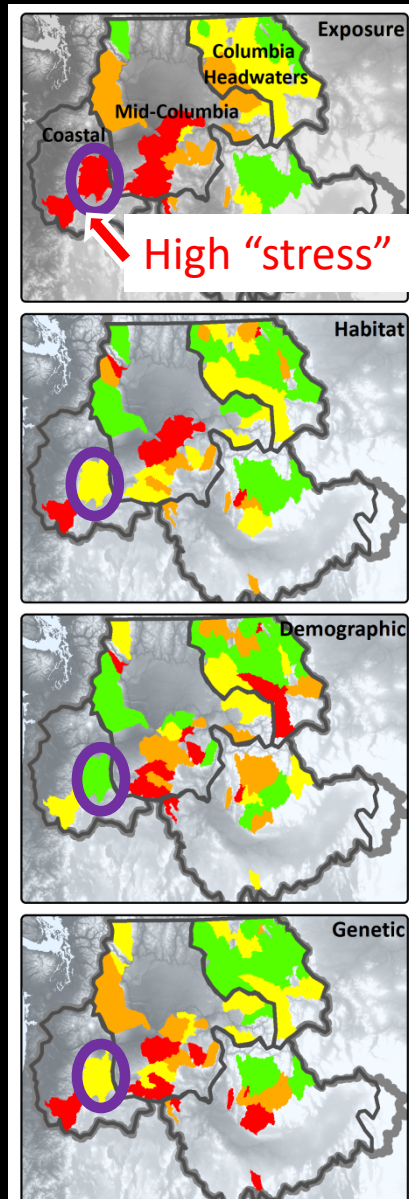
Demographics

Abundance
Life History Diversity
Productivity

Climate Change Vulnerability Assessment: How does demography and genetic factors interact with environmental factors across populations?

An example
from bull trout.
(Wade et al. 2016
with collaborators
from the USGS)

A given population
may have very
different levels of
“stress”
depending on
stressor type



A given population may be considered
“relatively vulnerable” solely on the
basis of the variables considered

How can we improve CCVAs?

Incorporating additional metrics into CCVAs and correlating with remote sensing data

Landscape Genetics/Genomics

Climate variables were correlated with genetic diversity in steelhead and bull trout genetic

(Hand et al. 2016; Kovach et al. 2015)



Demographic: productivity

Genetic: effective number of breeders (N_b)

Variation in Chinook population productivity is related to environmental conditions and habitat quality/quantity

Nick Gayeski – NGO: Wild Fish Conservancy (in prep)

Used Bayesian inference to correlate productivity with environmental variables.

Productivity
=Recruits/Adult Spawner
Long time series (~10-20 years) for 16 Chinook river basins

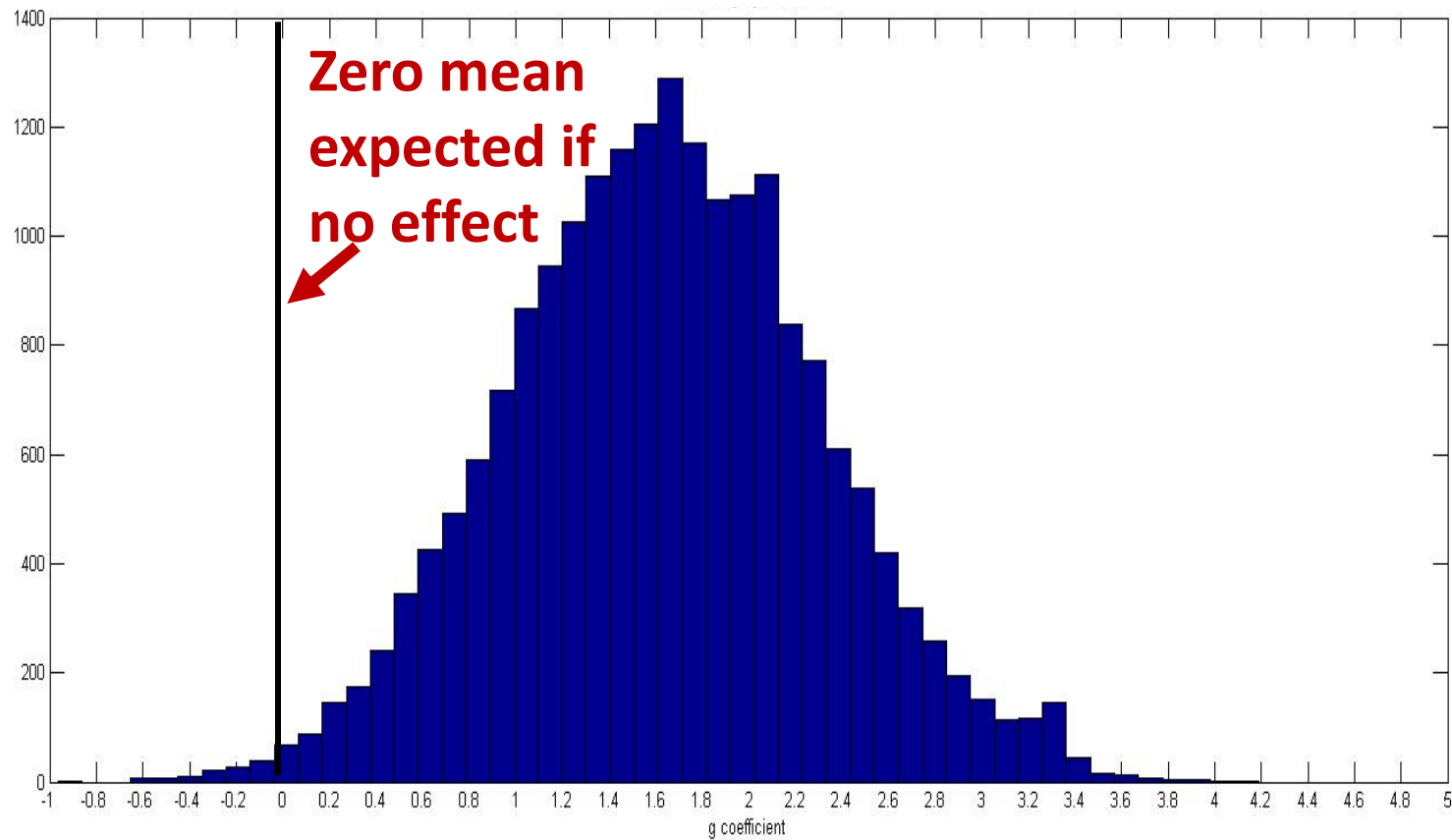
Environmental variables

- Stream temperature, floodplain area, the highest 5% winter flows, NPP, human disturbance



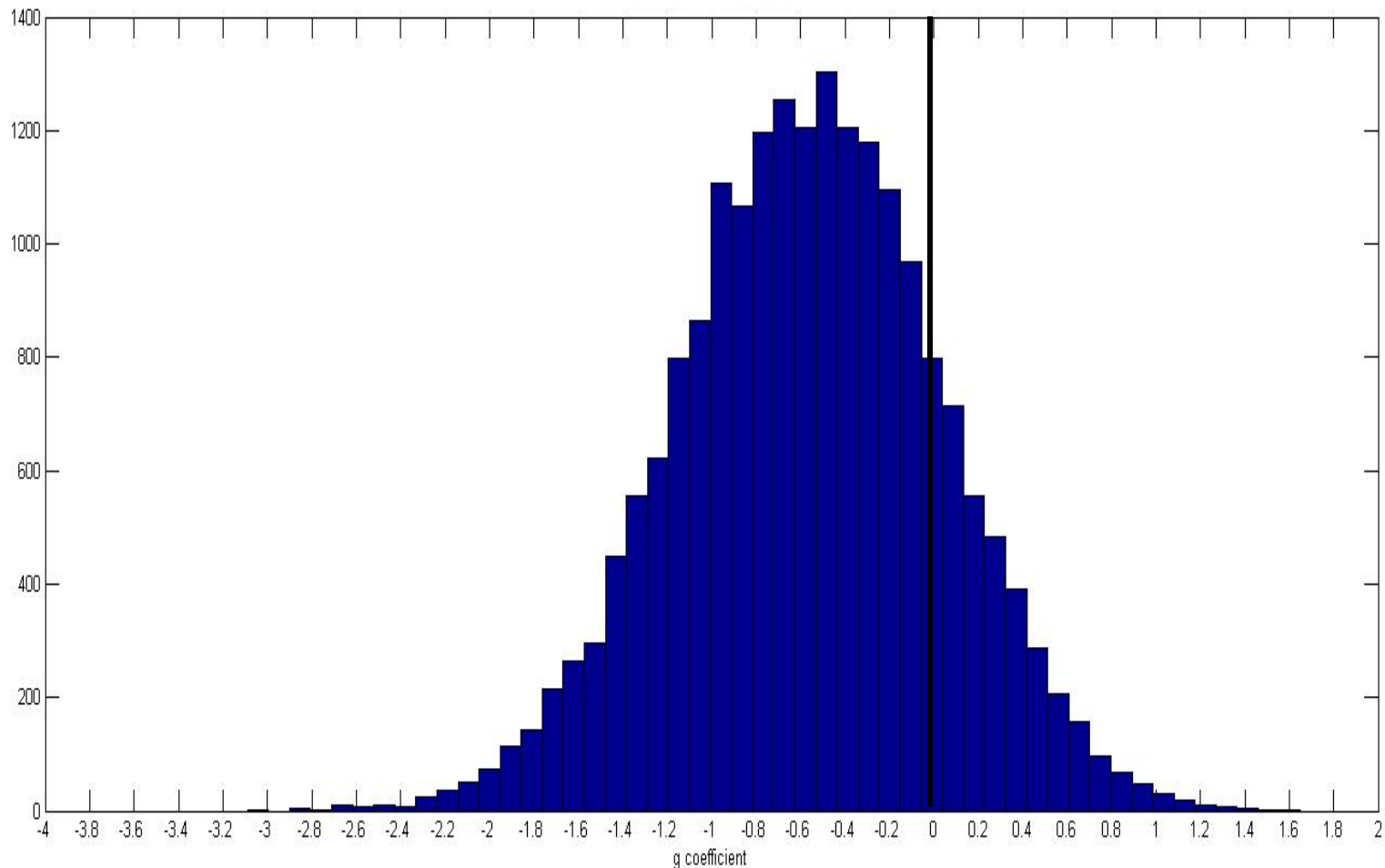
Chinook salmon population productivity (recruits/spawner) varies with stream temperature and NPP

Posterior Distribution of the Coefficient for Mean August Stream Temp (mean = 1.64)



Chinook salmon population productivity (recruits/spawner) varies with stream temperature and NPP

Posterior Distribution for the Coefficient for Mean NPP (mean = -0.57)



The Riverscape Analysis Project (RAP): An integrated platform for salmonid conservation



Web-based support tool for salmonid conservation

Basin scale data for the Pacific NW and fine scale (HUC 12) for the Columbia River Basin

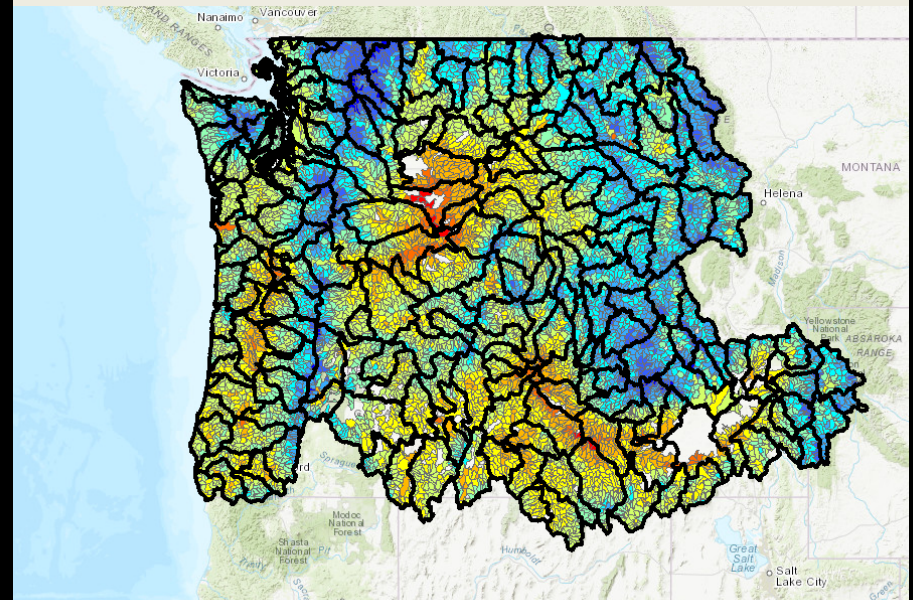
Data

- Expanded access to remotely-sensed climate/habitat data

Tools

- Basic CCVA
- Demogenetic monitoring (Nb)
- Landscape genetic tools

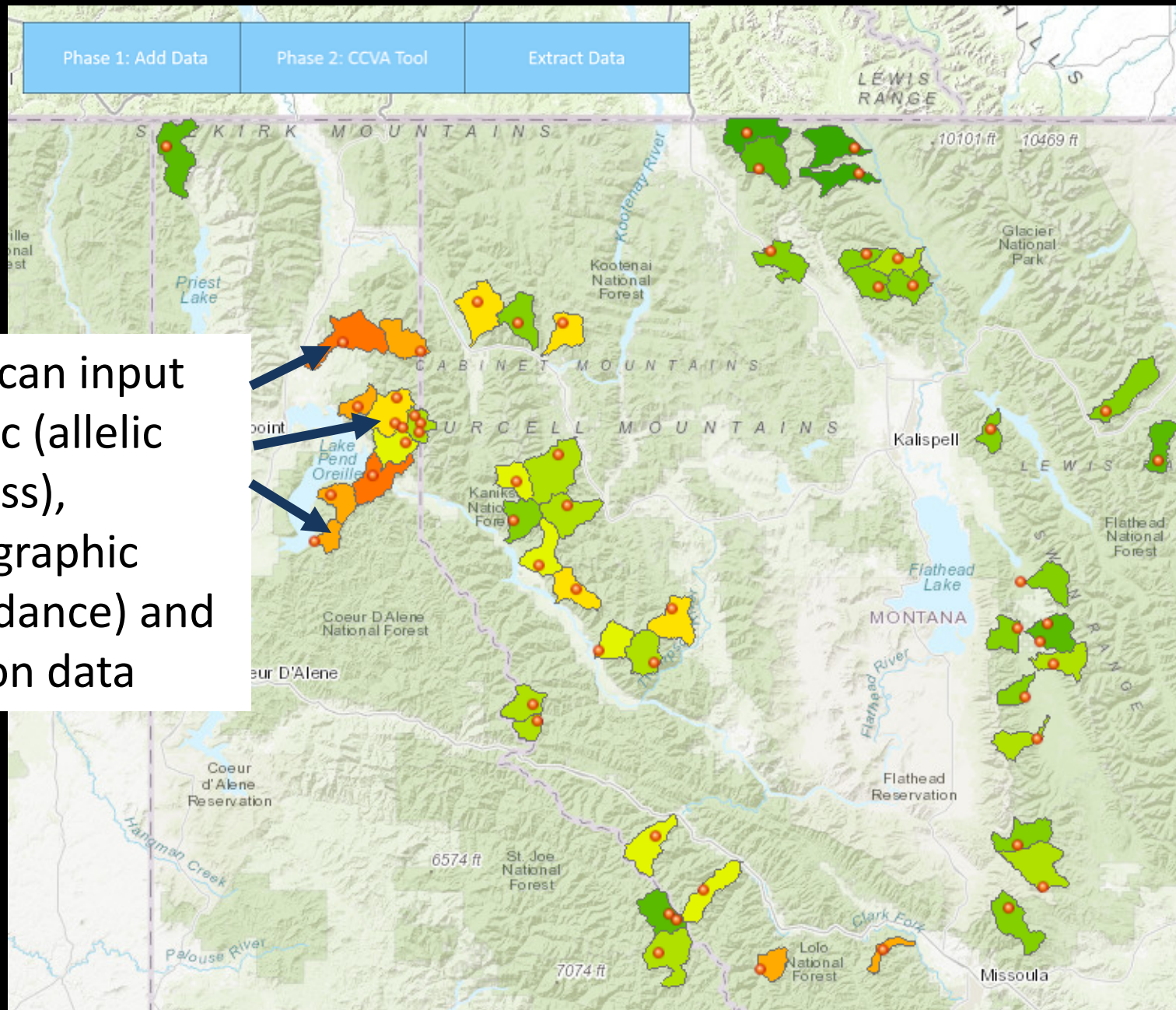
Columbia River Basin



Increased Data Access to salmonid/aquatic habitat characteristics

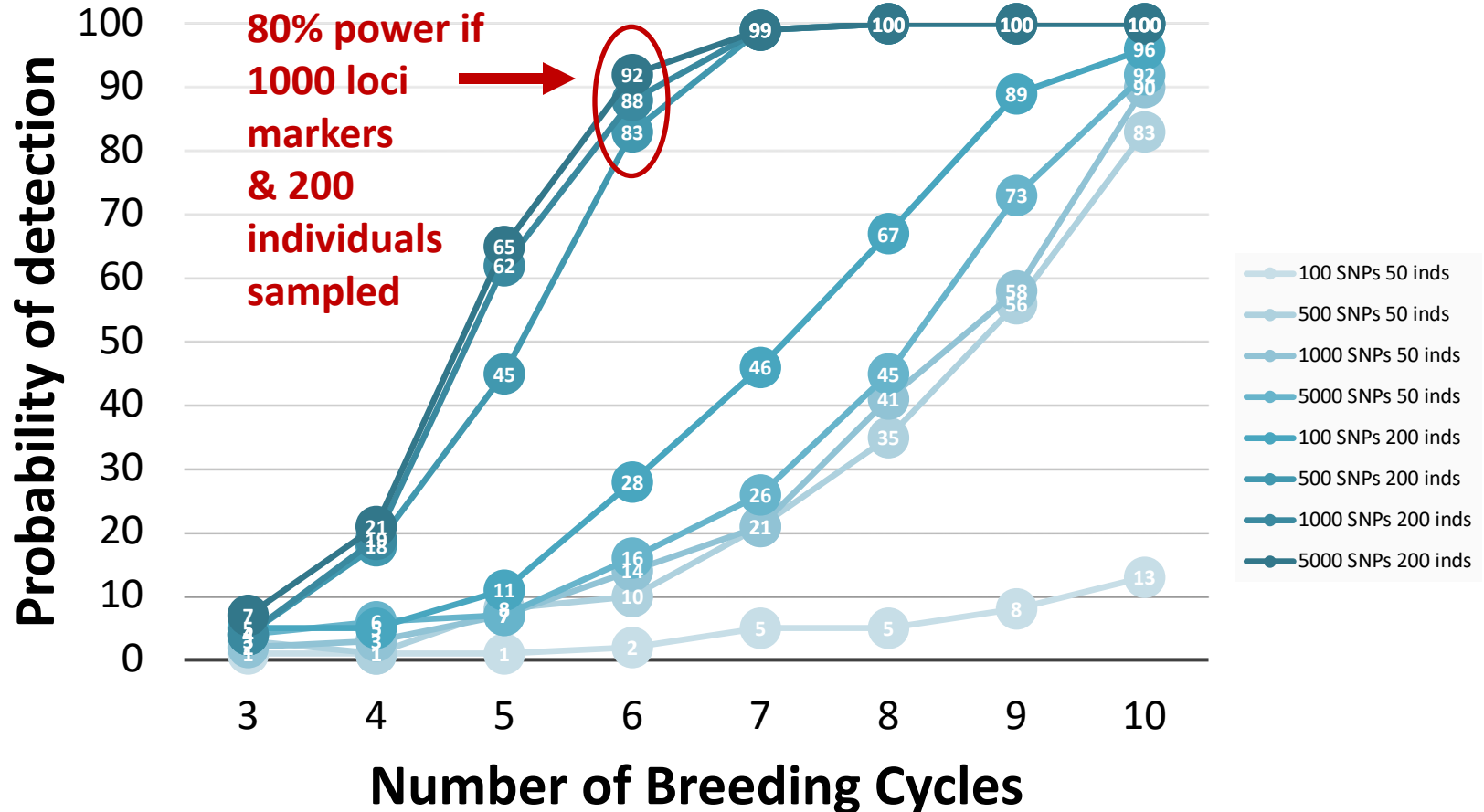
| Climate Data | RS Mission/ Product | Habitat Quality Data | RS Mission/ Product |
|--|---------------------------|--|---|
| Freeze-Thaw Timing | NASA SSM/I, AMSR-E | Drainage Density, Amount, and Sinuosity | NASA SRTM & NHDPlusv2 |
| Open Water | NASA AMSR-E | Productivity | NPP |
| NorWeST Stream Temperature | NASA Landsat TM & NAIP | Disturbance: NOAA CHAMP, Human Footprint, and NLCD 2011 | NASA GRUMP, GPWv3, DMSP, Landsat (Landcover - % disturbance and % forested) |
| USFS Stream Flow | n/a | Channel and Valley Slope | NASA SRTM |
| Future predictions Air Temperature, Precipitation, Runoff | NASA NEX-DCP 30 | Others: Glaciers, Dams, Elevation, Waterbodies | various |

Climate Change Vulnerability Assessment Tool Available now on RAP



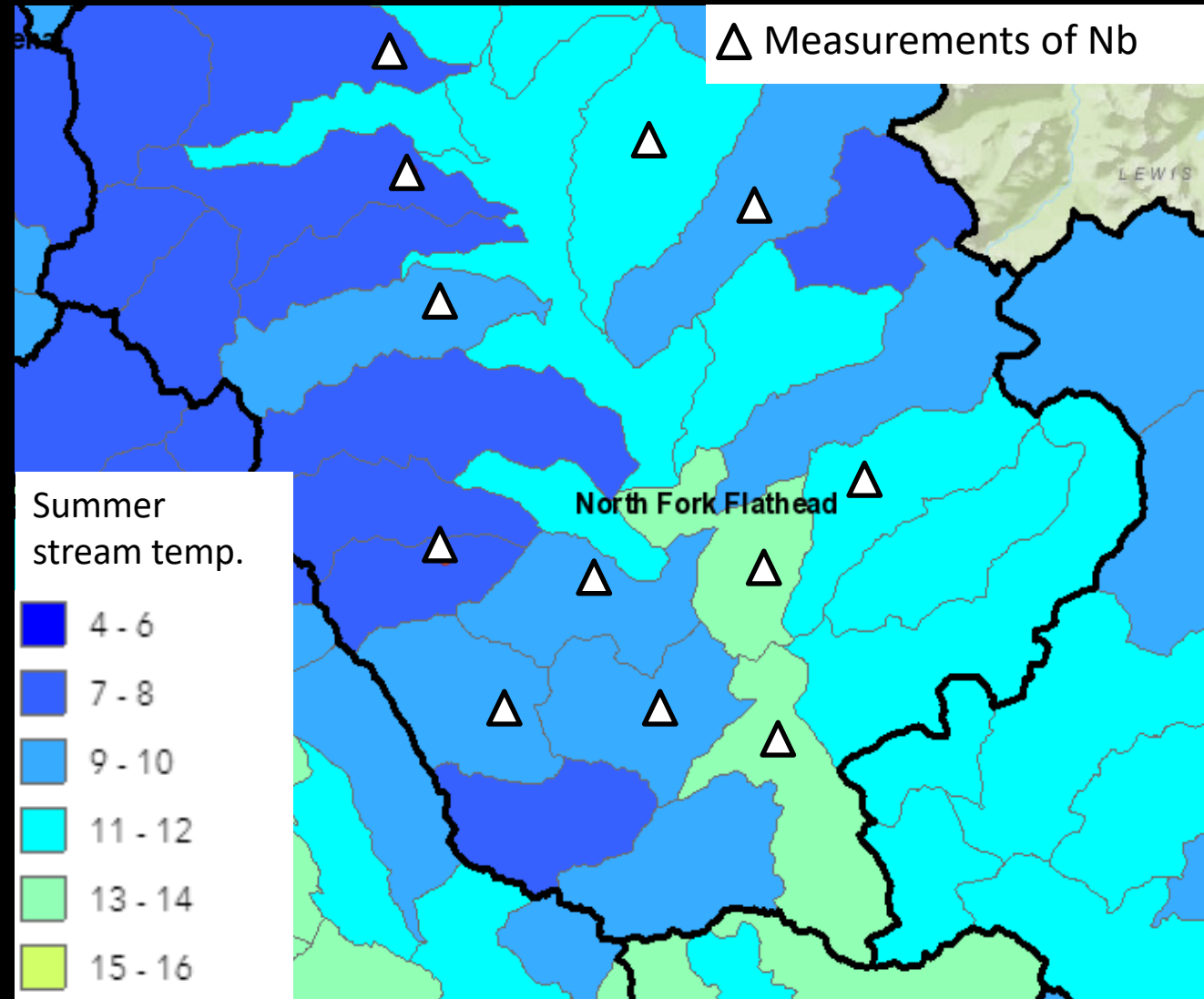
Answering the need for genomic monitoring tools related to abundance and genomic diversity: AgeStrucNe

Nb (number of breeders) provides an annual measure of abundance and genomic diversity, and is driven by environmental factors



Genomic Monitoring tools for resource managers: A tool for visualizing and correlating Nb to environmental variables

Coming Soon:
Available through
the RAP website



Advance training for project collaborators and conservation professionals

September 9-15th, 2018

CONGEN 2018

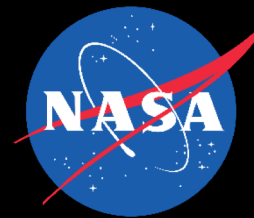
POPULATION GENOMICS
CONSERVATION GENETICS
COURSE

The course aims to provide training in conceptual and practical aspects of data analysis to understand the evolutionary and ecological genomics of natural and managed populations. Special section in this years course include landscape genomics, genomic monitoring tools for assessing the impacts of climate change and epigenomics.



For more information visit the website:
www.umontana.edu/sell/cps/congen2018/
Or contact Brian Hand or Gordon Luikart
Brian.Hand@umontana.edu
Gordon.Luikart@umontana.edu

Thanks!



Integrating remotely-sensed habitat quality and quantity, demographic, & genomic data

Integrating data, tools, & support (outreach) for salmonid conservation



pacific northwest aquatic
monitoring partnership



Montana Fish,
Wildlife & Parks



Columbia River
Inter-Tribal
Fish Commission

